

## SCD DNA Sequence (SEQ ID NO:1)

gtgggtgcgg	tgtcgccage	atccccggcg	ccctgctcg	gtcgccgag	ccctcgcc	60
ctgttctcct	ccccctcccg	cccttaccc	cacgcgggac	cgcccgcgc	agtcaactcc	120
tgcactttg	ccccgtcttg	gcagcggata	aaagggggt	gaggaaatac	cggacacgtc	180
cacccgttgc	cagctctagc	ctttaattc	ccggctcggg	acctccacgc	accgggctag	240
cggccacaac	cagctagcgt	gcaaggcggc	gccccgtcage	gctgtacccgc	gggttcgaa	300
accgcagtc	tccggcgacc	ccgaactccg	ctccggagcc	tcagccccct	gaaaagtgtat	360
cccgccatcg	gagagccaag	atgcggccc	acttgtcga	ggacgatatac	tctagctcct	420
ataccaccac	caccaccatt	acagcgcctc	cctccagggt	cctgcagaat	ggaggagata	480
agttggagac	gatgccc	tacttggaa	acgacat	ccctgtatata	aaagatgata	540
tatatgaccc	caccta	caag	gataagga	gcccaagccc	caaggttga	600
gaaacatcat	ccttatgtct	ctgctacact	tggagccct	gtatggatc	actttgattc	660
ctacctgcaa	gttctacacc	tggcttggg	gggtattcta	ctat	ttgtc	720
gcataaacagc	aggagtc	cgtctgtg	gcccacgc	ttacaaagct	cggctgcccc	780
ta	cggtctt	tctgtat	gccaacaca	ttggcattca	aatgtat	840
ctcg	tgacca	ccgtgcccac	cacaagttt	cagaacaca	tgctgat	900
gacgtggct	tttctctct	cacgtgggtt	ggctgtt	g	gcgcaaacac	960
aagagaaggg	gagtacg	gacttgc	acctaga	tgagaaactg	gtgtgttcc	1020
agaggaggt	ctacaaac	cttgc	tgatgt	catcctgc	acgcttgc	1080
cctgtatt	ctggggtgaa	actttcaaa	acagtgtt	cg	ttttgc	1140
atgctgtgt	gcttaat	acctggctt	tga	acagtgc	tgcccac	1200
gtccttat	ga	aaacatt	agccccggg	gaaat	ttcgat	1260
gtgagggctt	ccacaactac	caccact	ttcc	ctactctgc	agtgagtacc	1320
gctggcacat	caacttcacc	acattt	ttgat	ggccgc	ggtctggc	1380
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aaagcnagg	aaattgtcg	gggagag	tagcat	gaatgt	aggat	2040
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ccaccacata	gcatgttcc	tttctct	gg	aaa	atgg	2160
ggcaatgct	attcaat	gcaacatata	gtt	aggat	aaagat	2220
taagttt	gtaaaagtgg	tctgtc	gg	aggat	tttctt	2280
taacaagg	at	tcatat	ta	agg	tttctt	2340
tggtaaa	ac	agcagtc	agaa	ttt	caat	2400
ttctctt	ct	tc	tt	ttt	ttt	2460
ctgtggc	at	cc	cc	ttt	ttt	2520
aaataaaata	tatata	tata	tt	ttt	ttt	2580
ttccaaagag	ggatgtt	aaaaact	ga	ttt	ttt	2640
at	ct	ca	gg	ttt	ttt	2700
ttctaag	gaa	actt	gg	ttt	ttt	2760
tactg	gt	ttt	gg	ttt	ttt	2820
tccag	ctt	taaa	act	ttt	ttt	2880
ttcact	ttt	ttt	gg	ttt	ttt	2940
aactgc	ttt	ttt	gg	ttt	ttt	3000

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FIGURE 1

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2/24

taccctgtct	gtcccttttc	tttgaccaga	tctttcttctt	ccctgaacgt	tttcttcttt	3060
ccctggacag	gcagcctcct	ttgtgtgtat	tcagaggcag	tgtgacttg	ctgtccaggc	3120
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cttttctgt	attgggtggg	atttttccc	tttttatgt	ggatatagt	gtacttgt	5220
acaagaataa	ttttgaaata	atttctatta	atataactc	tgaagctaat	tgtactaata	5280
tgagattgt	tttggtcata	ataaaaagt	tgatgaatct	attgcact		5329

FIGURE 1 (CONT.)

3/24

## CA12 DNA Sequence (SEQ ID NO:2)

gtactcgcca	cggcacccag	gctgcgcgca	cgcgtcccc	gtgtcagct	ggagagcgag	60
cggccaccgg	gagcccccgg	cacagcccgc	gcccccggcc	caggagccg	cgaagatgcc	120
ccggcgacgc	ctgcacgcgg	cggccgtgt	cctgtggtg	atcttaaagg	aacagcccttc	180
cagccccggcc	ccagtgAACG	gttccaagt	gacttattt	ggtcctgtat	gggagaatag	240
ctggtccaag	aagtaccgt	cgtgtgggg	cctgtgcag	tcccccata	acctgcacag	300
tgacatccctc	cagtatgacg	ccagcctcac	gccctcgag	ttccaaggct	acaatctgtc	360
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cgagctgcac	attgtccatt	ataactcaga	cctttatcct	gacgcccagca	ctggcagcaa	600
caagtcaaaa	ggcctcgctg	tcctggctgt	tctcatttag	atgggctctt	tcaatccgtc	660
ctatgacaag	atcttcagtc	accttcaaca	tgtaaagtac	aaaggccagg	aagcattcgt	720
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caagccagcc	accaagatgg	agactgaggc	ccacgcttgc	ggtccccgg	gctccggggc	1200
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gacagcatga	aatgtctct	tgaagcatag	ctttttaaat	atcttttcc	ttctactct	2640
ccctctgact	ctaagaattt	tctcttctgg	aatcgcttgc	acccaggagg	cgagggttgc	2700
agtaagccaa	ggtcatgcca	ctgcactcta	gcctgggtga	cagagcgaga	ctccatctca	2760
aaaaaaaaaaa	aaaaaa					2775

FIGURE 2

4/24

## PIK3R4 DNA Sequence (SEQ ID NO:3)

gcacgagggg	agttcggcgt	ttgctgggc	tgcagcagct	gaagtgttagt	gttttcttgg	60
gactggcggt	ctgcacttct	ctcccccgggtt	ccatctcccc	ccgccccgggt	gtgaggccct	120
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FIGURE 3

acttggcagc	tttaggcata	actgggagac	aagtgtatct	tgttaaaacc	aaacaagaac	3060
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gcccagagtc	cctgcccgtg	ggacatcatg	acatcatc	tgtgtcgcc	acattcc	4560
ccacacagg	cttcatcgta	actgcttca	gagatgggat	tgtgaagg	tggaaataaaa	4620
acctactgt	ttgtataat	tttaatagtt	ataaaatataa	tactataact	cgagaaaagg	4680
catttctaga	gaacagattc	atttgc当地	ttttcaaaat	tatgtcttcc	tattactgtt	4740
tcatgactga	ctgactaaat	gacacccaaa	atggttaaga	tgtacttgac	tagttactt	4800
atgcata	ttgcaagaat	cagccagcc	acaatgtctg	ggattttat	tgtatatgtt	4860
atagaggtga	gaaatgtaaa	atatgaaaat	gaatatgtt	atttgtatt	aaaaagatg	4920
gttggaaaaga	tgggttgc当地	ctattatag	ataaaacacat	ttttgc当地	aaaaatgct	4980
ttcaaagcag	ttaaactgt	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaac	5040
tcgaggggggg	gccccgtacc					5060

FIGURE 3 (CONT.)

6/24

## PLD3 DNA Sequence (SEQ ID NO:4)

ctctttataa	tttagttcc	atagaagtta	tatgtgcatt	aaaaaaaaatt	caatgctgga	60
gcgaccgtgt	ctggggagcc	gagccccgct	tctcgctgcg	gtgagccgg	actggggcac	120
gcaactgcga	gactccccgc	tgcaagtggc	ggagtcccac	aggcccccc	cctcctccca	180
ccctcggtca	gcctgtccag	acagaagctg	gggcccagcg	gaggtagcag	cagacgcctg	240
agagcgagggc	cgaggccctc	agggtttgga	gaccctgaca	caccacctt	ctcacctggg	300
ctctcggtat	cccccagcct	tgagggaaaga	tgaagcctaa	actgtatgtac	cagagactga	360
aggtgcctgc	agaggagccc	gccaatgagc	tgcccatgaa	tgagatttag	gcgtggaagg	420
ctgcggaaaaaa	gaaagccgc	tgggtcctgc	tggtcctcat	tctggcggtt	gtgggcttcg	480
gaggcctgatg	actcaagctgt	ttctatggga	atacggcgac	ttgcacatct	ttgggccccaa	540
ccagcgccca	gccccctgct	atgacccttg	cgaagcagtg	ctggtgaaa	gcattcctga	600
gggcctggac	ttccccaaatg	cctccacggg	gaacccttcc	accagccagg	cctggctggg	660
cctgctcgcc	ggtgcgcaca	gcagcctgga	catgcctcc	ttctactgga	ccctcaccacaa	720
caatgacacc	cacacgcagg	agccctctgc	ccagcagggt	gaggaggccc	tccggcagct	780
gcagaccctg	gcacccaaagg	gcgtgaacgt	ccgcacatcgct	gtgagcaagc	ccagcggggcc	840
ccagccacag	gcggacactgc	aggctctgt	gcagagcgg	gcccagggtcc	gcatgggtgg	900
catgcagaag	ctgaccatg	gcgtcctgca	taccaagtcc	tgggtgggttgg	accagaccca	960
cttctacctg	ggcagtgcac	acatggactg	gcttcactg	acccaggta	aggagctggg	1020
cgtggtcatg	tacaactgca	gctgcctggc	tcgagacctg	accaagatct	ttgaggccta	1080
ctgggtcctg	ggccaggcag	gcagctccat	cccatcaact	tggcccccgtt	tctatgacac	1140
ccgctacaac	caagagacac	aatggagat	ctgcctcaat	ggaaccctg	ctctggccta	1200
cctggcgagt	gcgcccccac	ccctgtgtcc	aagtggccgc	actccagacc	tgaaggctct	1260
actcaacgtg	gtggacaatg	cccggagttt	catctacgtc	gctgtcatga	actacactgcc	1320
cactctggag	ttctcccacc	ctcacagggtt	ctggcctgcc	attgacgatg	ggctgcggcg	1380
ggccacctac	gagcgtggcg	tcaaggtgctg	cctgctcatac	agctgctggg	gacactcgga	1440
gccatccatg	cgggccttcc	tgctctctct	ggctgcctgc	cgtgacaacc	ataaccactc	1500
tgacatccag	gtgaaactct	ttgtggtccc	cgcggatgag	gcccaggctc	aatcccata	1560
tgcccgtgtc	aaccacaaca	agtacatgtt	gactgaacgc	gccacctaca	tcggaacctc	1620
caactggct	ggcaactact	tcacggagac	ggcggggcacc	tcgctgctgg	tgacgcagaa	1680
tgggagggggc	ggcctgcgga	gccagctgga	ggccattttc	ctgaggact	gggactcccc	1740
ttacattcat	gaccttgaca	cctcagctga	cagcgtggc	aacgcctgcc	gcctgctctg	1800
aggcccgatc	cagtggcag	gccaaggcct	gctggggcccc	cgcggaccca	ggtgctctgg	1860
gtcacggtcc	ctgtccccgc	accccccgtt	ctgtctgccc	cattgtggct	cctcaggctc	1920
tctccctgc	tctcccacct	ctacctccac	ccccaccggc	ctgacgctgt	ggcccccggga	1980
cccagcagag	ctggggggagg	gatcagcccc	caaagaaatg	ggggtgcatg	ctggcctgccc	2040
ccctggccca	cccccaactt	ccagggcaaa	aaggcccag	ggttataata	agtaaataac	2100
ttgtctgtaa	aaaaaaaaaaaa	aaaaaaaaaaaa	a			

FIGURE 4

7/24

## HSPD1 DNA Sequence (SEQ ID NO:5)

ggcacgaggc	gacgacctgt	ctcgccgagc	gcacgcctt	ccgccc	gcagaaatgc	60
ttcggttacc	cacagtctt	cgccagatga	gaccgtgtc	cagggta	cttc	120
tcactcgccc	ttatgccaaa	gatgtaaaat	ttgggtcaga	tgccc	gagcc	180
aagggttaga	ccttttagcc	gatgtgtgg	ccgttacaat	ggggccaa	gaaagaa	240
tgattattga	gcagagttgg	ggaagtccca	aagtaacaaa	agatgggtg	actgttgc	300
agtcaattga	cttaaaagat	aaatacaca	acattggagc	taaacttgtt	caagatgtt	360
ccaataaacac	aaatgaagaa	gctggggatg	gcactaccac	tgctactgt	ctggcac	420
ctatagccaa	ggaaggcttc	gagaagatta	gcaaagg	taatccag	gaaatcagg	480
gaggtgtgat	gttagctgtt	gatgtgtaa	ttgctga	actaa	ac	540
tgaccacccc	tgaagaaatt	gcacagg	ctacgat	tgcaaa	acgga	600
ttggcaaat	catctctgtat	gcaatgaaa	aagtgg	aaagggt	gtc	660
aggatggaaa	aacactgaat	gatgaatt	aaattatt	aggcat	gaa	720
gtatatattt	tccataactt	attaatac	caaaagg	gaaatgt	ttccagg	780
cctatgttct	gtttagt	gaa	aagaaaattt	ctagtat	gtccatt	840
aaattgccaa	tgctcaccgt	aagccttgg	tcataat	tgc	gatgg	900
ctctaagtac	actcg	tta	aggctt	tcagg	gttgc	960
ctccagggtt	ttgtgaca	aat	agctt	tatgg	actgg	1020
gtgcagtgtt	tggagaag	ggat	gaccc	tgaatctt	agacgtt	1080
taggaaaagt	tggagagg	tc	attgtgac	aaagacgat	catgtctt	1140
gtgacaaggc	tcaaattgaa	aaacgtat	tt	agg	gttgc	1200
ctagtgaata	tgaaaaggaa	aaactgaa	tttgc	aaaactt	gatgg	1260
ctgtgctgaa	ggttgg	ttgg	acaagtgt	ttgaa	gacag	1320
cagatgc	ctt	ttgg	ttgtgt	ttgaa	tttgc	1380
gtgc	tcgat	ccag	ttgg	actcatt	ttccag	1440
aaattggat	agaaattt	aaaaga	tttgc	ttca	tttgc	1500
atgcagg	tgaagg	tttgc	ttgtgt	tttgc	tttgc	1560
gttatgt	tatgg	ttttgt	ttatgt	tttgc	tttgc	1620
caaagg	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	1680
cagaagg	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	1740
gtgg	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	1800
ttat	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	1860
gtc	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	1920
gtt	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	1980
ttat	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	2040
tgt	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	2100
aatcagg	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	2160
gtgagaata	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	2220
aaaaattt	ttatgg	ttatgg	ttgtgt	tttgc	tttgc	2258

FIGURE 5

8/24

## ZPK Variant 2 DNA Sequence (SEQ ID NO:6)

agcatccccga	gcccggagctgc	agcagcgccg	ccttttgc	tgccggccgc	gagcccccca	160
gggccccagtgt	ttcacccatca	taccaggggc	cagaggcgat	ggcttcgc	catgagaccc	120
gaacacccctc	tccttcctt	gggggccttgc	tgtctaccct	aagtgaggca	tccatgcgc	180
agctggaccc	agacacttct	gactgcactc	ccgagaagga	cctgacgcct	accatgtcc	240
tgcagctaca	tgacgaggat	gcagggggcc	cagggggagc	agctgggtca	cctgagagtc	300
gggcattccag	attcgagct	gacgaggtgc	gactgcagtg	ccagagtggc	agtggcttcc	360
ttgagggcct	cttggctgc	ctgcgccttgc	tctggaccat	gattggcaaa	gcctactcca	420
ctgagcacaa	gcagcagcag	gaagaccttgc	gggaggtccc	ctttgaggaa	atcctggacc	480
tgcagtgggt	gggctcaggg	gcccaggggt	ctgtcttcc	ggggcgcttc	cacggggagg	540
aggtgtgtgt	gaagaagggt	cgagacctca	aagaaaaccga	catcaagcac	ttgcgaaagc	600
tgaagcaccc	caacatcatc	actttcaagg	gtgtgtgcac	ccaggctccc	tgctactgca	660
tcctcatgga	gttctgcgcc	cagggccagc	tgtatgagggt	actgcgggct	ggccgcctgc	720
tcacccctc	cttactgggt	gactggtcca	tgggcacatgc	tggtgcatg	aactacctgc	780
acctgcacaa	gattatccac	agggatctca	agtcacccaa	catgctaatac	acctacgacg	840
atgtggtgaa	gatctcagat	tttggacttgc	ccaaggagct	gagtgacaag	agcaccaaga	900
tgtcccttgc	agggacagta	gcctggatgg	cccctgaggt	gatccgcaat	gaacctgtgt	960
ctgagaaggt	cgacatctgg	tcctttggcg	ttgtgtatgc	ggaactgctg	actgggtgaga	1020
tccctacaa	agacgttagat	tcctcagcca	ttatctgggg	tgtgggaagc	aacagtctcc	1080
atctgcccgt	gcctccagat	tgcccagatg	gtttcaagat	cctgcttcgc	cagtgtcgga	1140
atagcaaacc	acgaaatcgc	ccatcattcc	gacagatctc	gctgcatctg	gacattgcct	1200
cagctgatgt	actctccaca	ccccaggaga	cttactttaa	gtcccaggca	gagtggcg	1260
aagaagtaaa	actgcacttt	gaaaagatta	agtcagaagg	gacctgtctg	caccgcctag	1320
aagaggaact	ggtgatgagg	aggagggagg	agctcagaca	cgccctggac	atcagggagc	1380
actatgaaag	gaagctggag	agagccaaca	acctgtatata	ggaacttaat	gccctcatgt	1440
tgcagctgga	actcaaggag	agggagctgc	tcagggcgaga	gcaagcttta	gagcggaggt	1500
gcccagggcct	gctgaagcca	cacccttccc	ggggcctcct	gcatggaaac	acaatggaga	1560
agcttatcaa	gaagaggaat	gtgccacaga	agctgtcacc	ccatagaaaa	aggccagata	1620
tcctcaagac	ggagtcttgc	ctccctaaac	tagatgcagc	cctgagtgccc	gtggggcttc	1680
ctgggtgtcc	taaggcccc	ccctcaccag	gacggagtcg	ccgtggcaag	accgcgtcacc	1740
gcaaggccag	cgccaagggg	agctgtgggg	acctgcctgg	gcttcgtaca	gctgtgccac	1800
cccatgaacc	tgaggagcca	ggaagccag	ggggccttagg	agggggaccc	tcagcctggg	1860
aggcctgccc	tccgcctc	cgtgggttc	atcatgacct	cctgctccgc	aaaatgtctt	1920
catcgcccc	agacctgtc	tcagcagcac	tagggtcccc	gggccccggg	gccacaggcg	1980
gagctgggaa	tcttgcctca	ccacccccc	ccgggggtga	cacccacca	agtgagggct	2040
cagcccttgc	ctccaccagc	ccagattcac	ctgggggagc	caaaggggaa	ccacccctcc	2100
cagtagggcc	tgttgaagggt	gtggggcttc	tggaaactgg	aagggaaagg	acctcaggcc	2160
gggggaggaag	ccgggctggg	tcccaact	tgacccca	tgcactgtc	tacagggctg	2220
ccgtcaccc	aagtcaaaaa	cgtggcatct	catcggaaaga	ggagggaaagg	gaggttagaca	2280
gtgaagtaga	gctgacatca	agccagaggt	ggcctcagag	cctgaacatg	cgccagtcac	2340
tatctacctt	cagctcagag	aatccatcag	atggggagga	aggcacagct	agtgaaccc	2400
cccccaagtgg	cacacctgaa	gttggcagca	ccaacactg	tgagggc	gatgagcggt	2460
ctgatgacat	gtgctccca	ggctcagaaa	tcccaactgg	cccacccct	tcagaggtca	2520
tccctggccc	tgaaccc	tccctgccc	ttccacacca	ggaacttctc	agagagcg	2580
gcccctccaa	ttctgaggac	tcagactgt	acagcactga	attggacaac	tccaaacagc	2640
ttgatgcctt	ggggccccca	gcttccctc	ctccatgaaa	gccactcgta	ttcttgta	2700
atagagaaat	atttatatgg	atttatata	tatacatata	tatatatata	tgcccccacat	2760
aatcaacaga	aagatggggc	tgtccca	gtaagtca	ctcgaggag	actgatcccc	2820
tgaccaattc	acctgataaa	ctctagg	actggcagct	gtggaaatga	atgaggcaca	2880
gcccgtagagc	tgtggctaa	ggcaagcccc	ttcctgcccc	acccattcc	ttatattc	2940
caagcaacaa	ggcaatagaa	aagccagggt	tgtcttata	ttcttata	ccaaataata	3000
gggggtgggg	ggagggcg	tgggggggc	aggagagaaa	accacttaga	ctgcacttt	3060
ctgttcgtt	tactctgtt	acacatttt	cacttggag	gagggaggct	aaggctgggt	3120
cctccctct	gaggttctc	aggtggcaat	gtaactcatt	tttttgc	accatttac	3180
ttctctgccc	aagccctgtc	ttaaggccca	gggggaggt	aggagactga	tagcatgtga	3240
tggctcaggc	tgaagaaccg	gggtgtgtt	taagtccctg	cttttatcct	ggtgcctgtat	3300
ttgggtgggg	actgtcctac	tgtaa	gtaaaaacc	ttgaaatata	acactccatg	3360
caagga						3365

## FIGURE 6

9/24

## SCD Amino Acid Sequence (SEQ ID NO:7)

MPAHLQLDDI	SSSYTTTTTI	TAPPSRVLQN	GGDKLETMPL	YLEDDIRPDI	KDDIYDPTYK	60
DKEGPSPKVE	YVWRNIIILMS	LLHLGALYGI	TLIPTCKFYT	WLWGVFYFYFV	SALGITAGAH	120
RLWSHRSYKA	RLPLRLFLII	ANTMAFQNDV	YEWARDHRAH	HKFSETHADP	HNSRRGFFFS	180
HVGWLLVRKH	PAVKEKGSTL	DLSDEAEKL	VMFQRRYYKP	GLLLMCFILP	TLVPWYFWGE	240
TFQNSVFVAT	FLRYAVVLNA	TWLVNSAAHL	FGYRPYDKNI	SPRENILVSL	GAVGEGFHNY	300
HHSFPYDYS	SEYRWHINFT	TFFIGDCMAAL	GLAYDRKKVS	KAAILARIKR	TGDGNYKSG	359

FIGURE 7

## CA12 Amino Acid Sequence (SEQ ID NO:8)

MPPRSSLHAAA	VLLLWILKEQ	PSSPAPVNNGS	KWTYFGPDGE	NSWSKKYPSC	GGLLQSPIDL	60
HSDILQYDAS	LTPLEFQGYN	LSANKQFLLT	NNGHHSVKLNL	PSDMHIQGLQ	SRYSATQLHL	120
HWGNPNDPHG	SEHTVSGQHF	AAELHIVHYN	SDLYPDASTA	SNKSEGLAVL	AVLIEMGSFN	180
PSYDKIFSHL	QHVKYKGQEA	FVPGFNIEEL	LPERTAEYYR	YRGSLTPPC	NPTVLWTVFR	240
NPVQISQEQL	LALETALYCT	HMDDPSPREM	INNFRQVQKF	DERLVYTSFS	QVQVCTAAGL	300
SLGIILSLAL	AGILGICIVV	VVSIWLFRKK	SIKKGDNKGV	IYKPATKMET	EAHA	354

FIGURE 8

## PIK3R4 Amino Acid Sequence (SEQ ID NO:9)

MGNQLAGIAP	SQILSVESYF	SDIHDFEYDK	SLGSTRFFKV	ARAKHREGLV	VVKVFAIQDP	60
TLPLTSYKQE	LEELKIRLNS	AQNCLPFQKA	SEKASEKAAM	LFRQYVRDNL	YDRISTRPFL	120
NNIEKRWIAF	QILTAVDQAH	KSGVRHGDIK	TENVMTSWN	WVLLTDFASF	KPTYLPEDNP	180
ADFNYFFDTS	RRRTCYIAPE	RFVDGGMFAT	ELEYMRDPST	PLVDLNSNQR	TRGELKRAMD	240
IFSAAGCIAE	LFTEGVPLFD	LSQLLAYRNG	HFFPEQVLNK	IEDHSIRELV	TQMIHREPDK	300
RLEAEEDYLKQ	QRGNAFPEIF	YTFLQPYMAQ	FAKETFLSAD	ERILVIRKDL	GNIIHNLCGH	360
DLPEKAEGEP	KENGLVILVS	VITSCLQTLK	YCDSKLAALE	LILHLAPRLS	VEIILDRITP	420
YLLHFSNDSV	PRVRAEALRT	LTKVLALVKE	VPRNDINIYP	EYILPGIAHL	AQDDATIVRL	480
AYAENIALLA	ETALRFLELV	QLKNLNFMEND	PNNEEIDEVT	HPNGNYDTEL	QALHEMVQQK	540
VVTLLSDPEN	IVKQTLMENG	ITRLCVFFGR	QKANDVLLSH	MITFLNDKND	WHLRGAFFDS	600
IVGVAAYVGW	QSSSIKPLL	QQGLSDAEEF	VIVKALYALT	CMCQLGLLQK	PHVYEFASDI	660
APFLCHPNLW	IRYGAVGFIT	VVARQISTAD	VYCKLMPYLD	PYITQPIIQL	ERKLVLLSVL	720
KEPVRSRISFD	YALRSKDITS	LFRHLHMRQK	KRNGLSLPDCP	PPEDPAIAQL	LKKLLSQGMT	780
EEEEEDKLLAL	KDFMMKSNKA	KANIVDQSHL	HDSSQKGVID	LAALGITGRQ	VDLVTKQEP	840
DDKRARKHVK	QDSNVNEEWK	SMFGSLDPPN	MPQALPKGSD	QEVIQTGKPP	RSESSAGICV	900
PLSTSSQVPE	VTTVQNKPKV	IPVLSSTILP	STYQIRITTC	KTELQQLIQQ	KREQCNAERI	960
AKQMMENAEW	ESKPPPPGWR	PKGLLVAHLH	EHKSAVNRI	VSDEHSLFAT	CSNDGTVKI	1020
NSQKMEGKTT	TTRSILTYSR	IGGRVKTLLF	CQGSHYLAIA	SDNGAVQLLG	IEASKLPKSP	1080
KIHPLQSRIL	DQKEDGCVVD	MHHFNNSGAQS	VLAYATVNGS	LGVWDLRSSS	NAWTLKHDLK	1140
SGLITSFAVD	IHQCWLCIGT	SSGTMACWDM	RFQLPISSH	HPSRARIRRL	SMHPLYQSWV	1200
IAAVQGNNEV	SMWDMETGDR	RFTLWASSAP	PLSELQPSPH	SVHGIYCSPA	DGNPILLTAG	1260
SDMKIRFWDL	AYPERSYVVA	GSTSSPSVSY	YRKIIEGTEV	VQEIQNKQKV	GPSDDTPRRG	1320
PESLPVGHH	IITDVATFQT	TQGFIVTASR	DGIVKVWK			1358

FIGURE 9

10/24

## PLD3 Amino Acid Sequence (SEQ ID NO:10)

MTQLFLWEYG DLHLFGPNQR PAPCYDPCEA VLVESIPEGL DFPNASTGNP STSQAWLGLL	60
AGAHSSLDIA SFYWTLTNNND THTQEPSAQG GEEVLRQLQT LAPKGVNVRV AVSKPSGPQP	120
QADLQALLQS GAQVRMVDMQ KLTHGVLHTK FWVVVDQTHFY LGSANMDWRS LTQVKELGVV	180
MYNCSCLARD LTKIFEAYWF LGQAGSSIPS TWPRFYDTRY NQETPMEICL NGTPALAYLA	240
SAPPPLCPSG RTPDLKALLN VVDNARSFIY VAVMNYLPTL EFSHPHRFWP AIDDGLRRAT	300
YERGVKVRLR ISCWHGSEPS MRAFLLSLAA LRDNHHTHSDI QVKLFVVPAD EAQARI PYAR	360
VNHNKYMVT RATYIGTSNW SGNYFTETAG TSLLVTQNGR GGLRSQLEAI FLRDWDSPI	420
HDLDTSADSV GNACRLL	437

FIGURE 10

## HSPD1 Amino Acid Sequence (SEQ ID NO:11)

MLRLPTVFRQ MRPVSRVLAP HLTRAYAKDV KFGADARALM LQGVDLLADA VAVTMGPKG	60
TVIIEQSWGS PKVTKDGVTW AKSIDLKDKY KNIGAKLVQD VANNTNEEAG DGTTTATVLA	120
RSIAKEGFEK ISKGANPVEI RRGVMLAVDA VIAELKKQSK PVTTPEEIAQ VATISANGDK	180
EIGNIISDAM KKVGKGVIT VKDGKTLNDE LEIIIEGMKFD RGYISPYFIN TSKGQKCEFQ	240
DAYVLLSEKK ISSIQSIVPA LEIANAHRKPV LVIIAEDVDG EALSTLVLNR LKVGLQVVAV	300
KAPGFGDNRK NQLKDMAIAT GGAVFGEEGL TLNLEDVQPH DLGKVGEIVV TKDDAMLLKG	360
KGDKAQIEKR IQEIIIEQLDV TTSEYEKEKL NERLAKLSDG VAVLKVGCTS DVEVNEKKDR	420
VTDALNATRA AVEEGIVLGG GCALLRCIPA LDSLTPANED QKIGIEIIR TLKIPAMTIA	480
KNAGVEGSLI VEKIMQSSSE VGYDAMAGDF VNMVEKGIID PTKVVRTALL DAAGVASLLT	540
TAEVVVTEIP KEEKDPMGA MMMGGGGMGF GMF	573

FIGURE 11

## ZPK Variant 2 Amino Acid Sequence (SEQ ID NO:12)

MACLHETRTP SPSFGGFVST LSEASMRKLD PDTSDCTPEK DLTPTHVLQL HEQDAGGPGG	60
AAGSPESRAS RVRADEVRLQ CQSGSGFLEG LFGCLRPVWT MIGKAYSTEH KQQQEDLWEV	120
PFEEILDLQW VGSGAQGAVF LGRFHGEEVA VKKVRDLKET DIKHLRKLKH PNIITFKGVC	180
TQAPCYCILM EFCAQGQOLYE VLRAAGRPPVT SLLVWDWSMGI AGGMNYLHLH KIIHDLKSP	240
NMLITYDDVV KISDFGTSKE LSDKSTKMSF AGTVAWMAPE VIRNEPVSEK VDIWSFGVVL	300
WELLTGIEPY KDVDSSAIIW GVGSNSLHLH VPSSCPDGFK ILLRQCWN SK PRNRPSFRQI	360
LLHLDIASAD VLSTPQETYF KSQAEWREEV KLHFKEKIKSE GTCLHRLEEE LVMRRREELR	420
HALDIREHYE RKLERANNLY MELNALMQL ELKERELLRR EQALERRCPG LLKPHPSRGL	480
LHGNTMEKLI KKRNPQKLS PHSKRPDIK TESLLPKLDA ALSGVGLPGC PKAPPSPGRS	540
RRGKTRHRKA SAKGSCGDLP GLRTAVPPHE PGPGPGSPGGL GGGPSAWEAC PPALRGLHHD	600
LLLRKMSSSS PDLLSAALGS RGRGATGGAG DPGSPPPPARG DTPPSEGSAP GSTSPDSPGG	660
AKGEPPPPVG PGEVGVL LGT GREGTSGRGG SRAGSQHLTP AALLYRAAVT RSQKRGISSE	720
EEEVEVDSEV ELTSSQRWPQ SLNMRQSLST FSSENPSDGE EGTASEPSPS GTPEVGSTNT	780
DERPDERSDD MCSQGSEIPL DPPPSEVIPG PEPSSLPIPH QELLRERGPP NSESDCDCST	840
ELDNSNSVDA LRPPASLPP	859

FIGURE 12

11/24

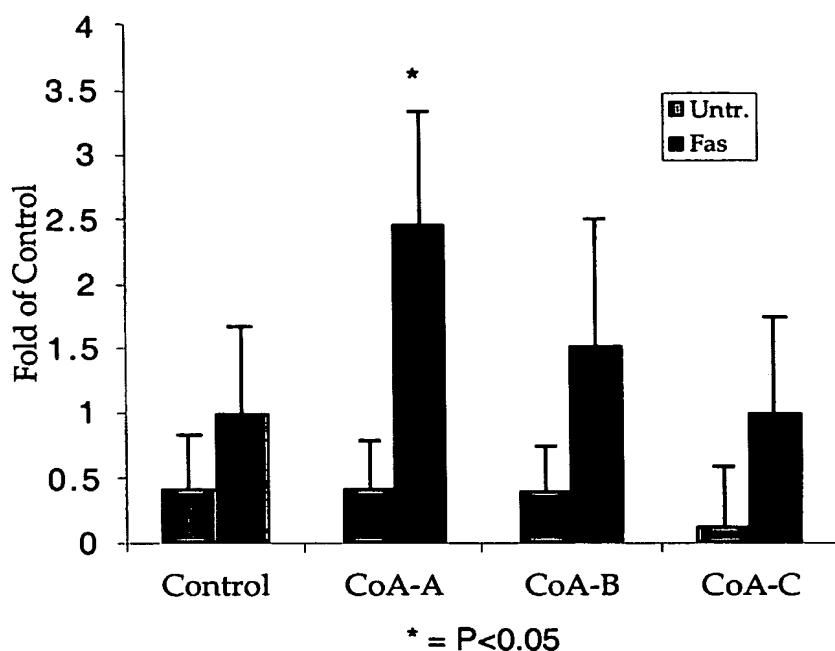


FIGURE 13A

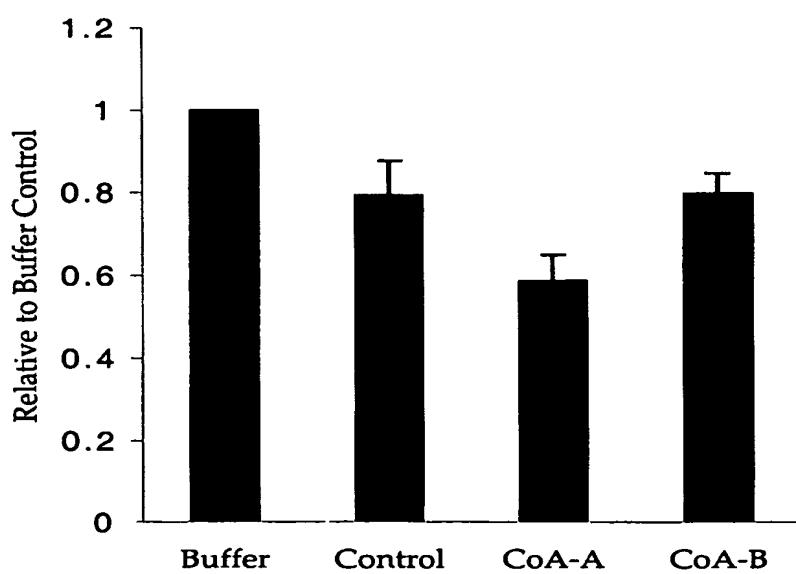


FIGURE 13B

12/24

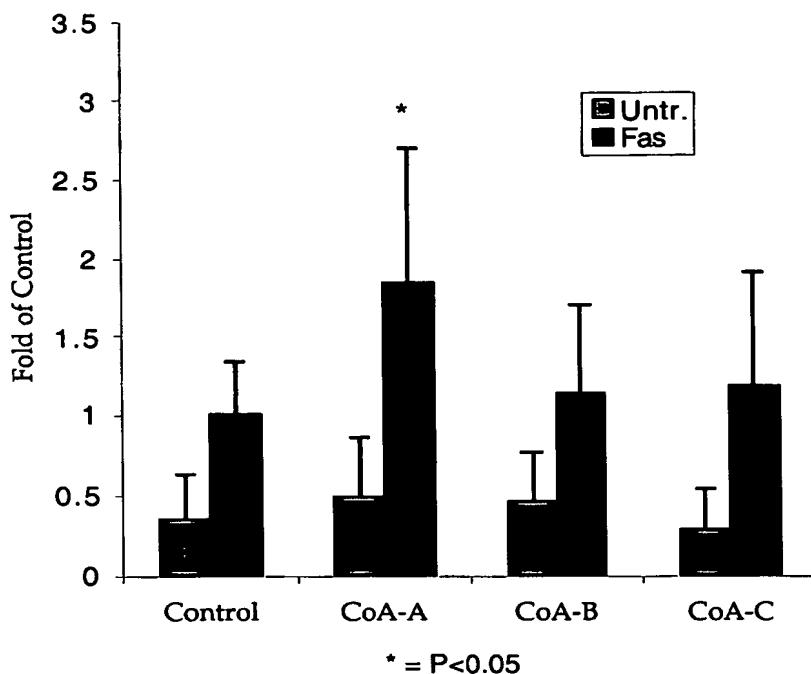


FIGURE 14A

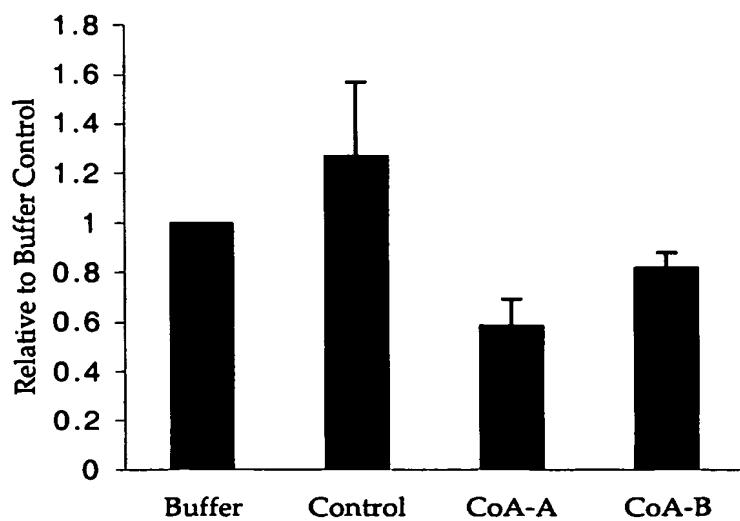


FIGURE 14B

13/24

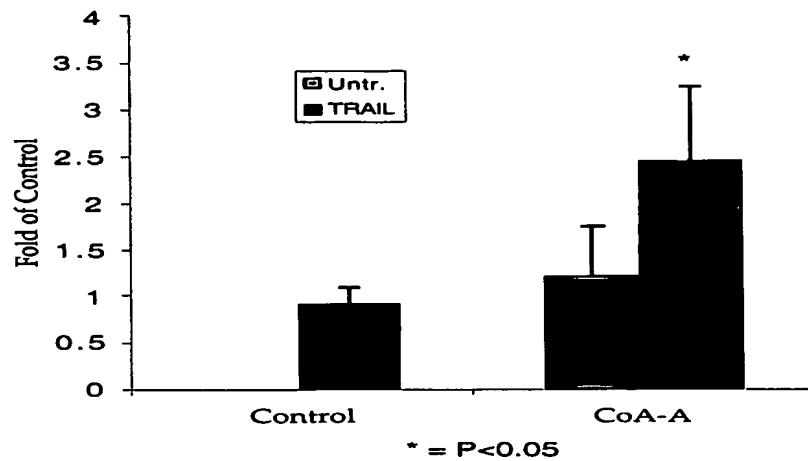


FIGURE 15A

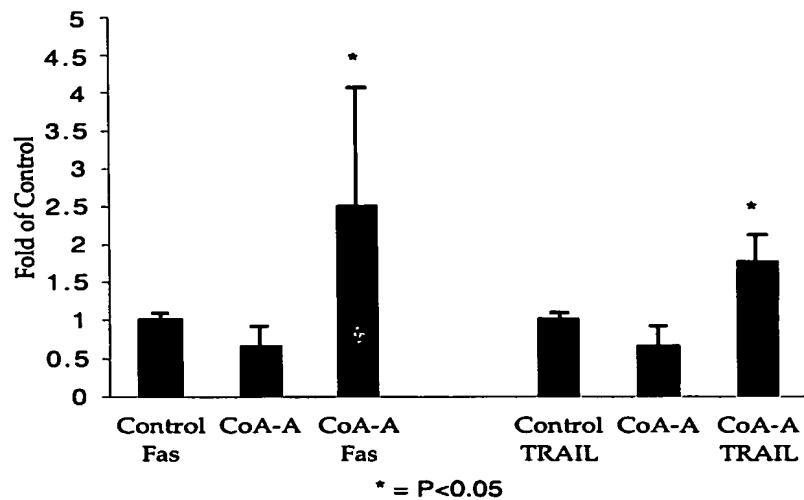


FIGURE 15B

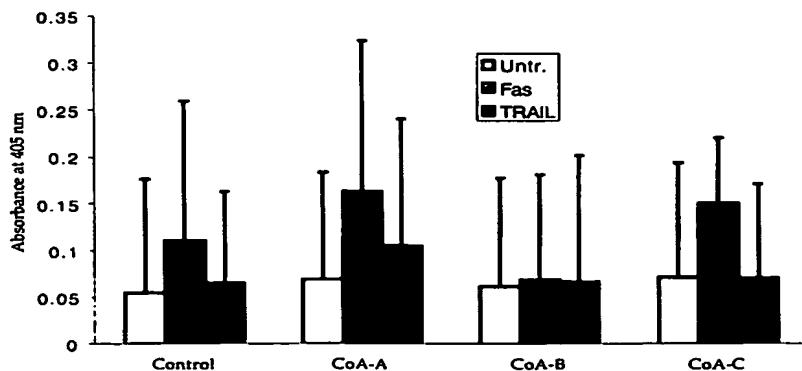


FIGURE 16

14/24

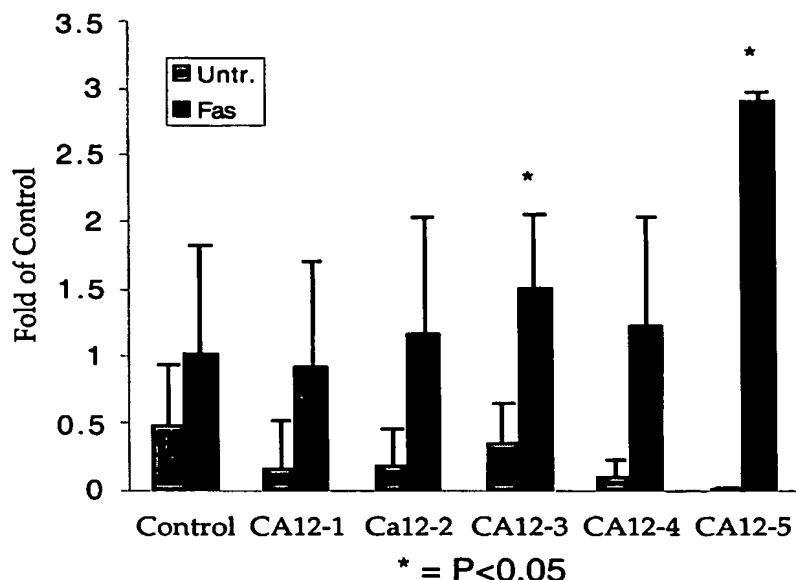


FIGURE 17A

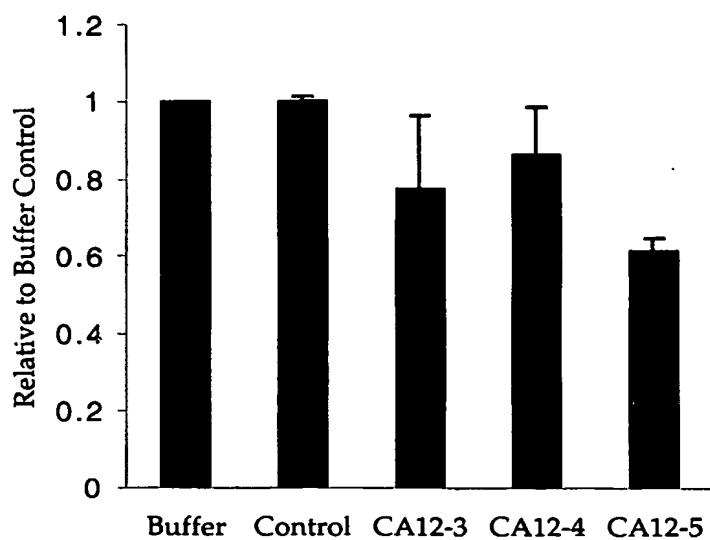


FIGURE 17B

15/24

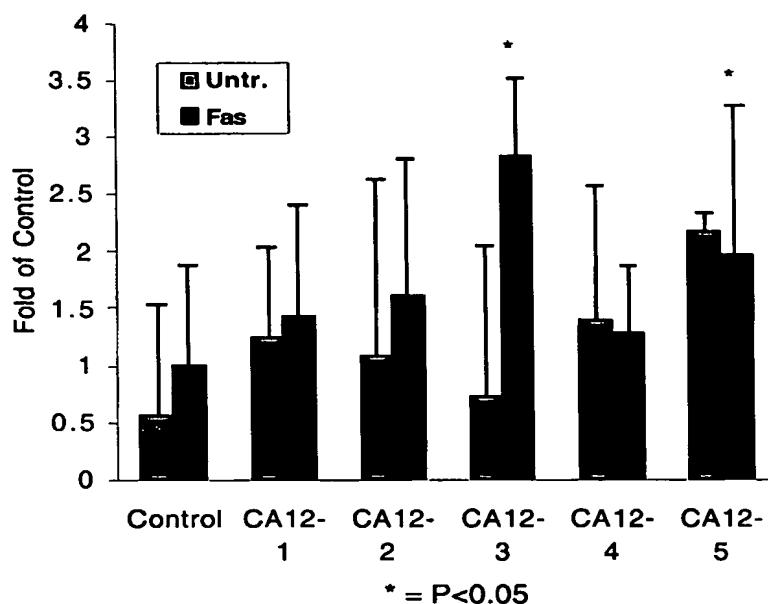


FIGURE 18A

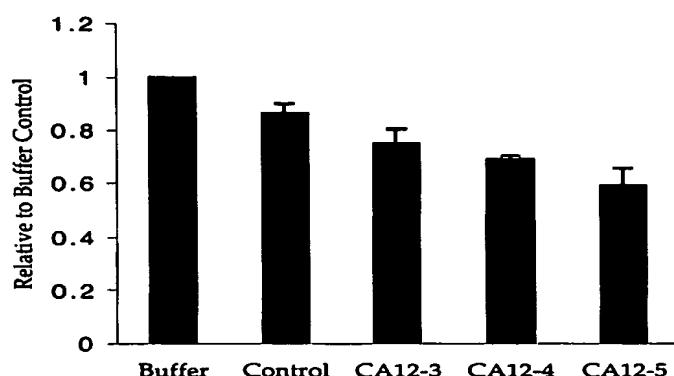


FIGURE 18B

16/24

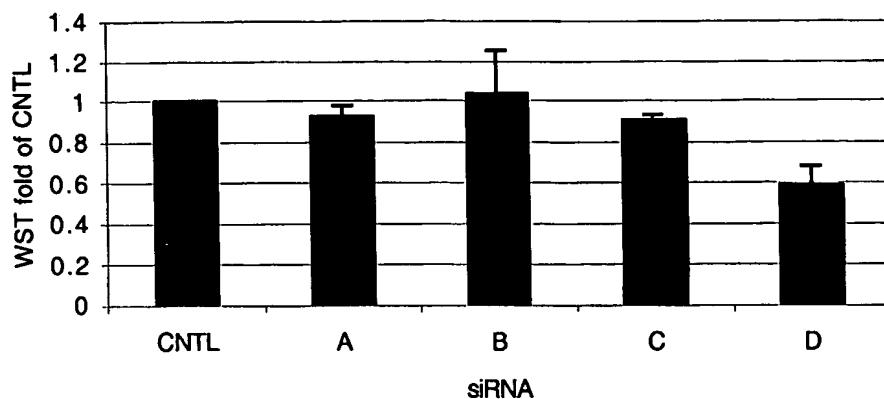


FIGURE 19

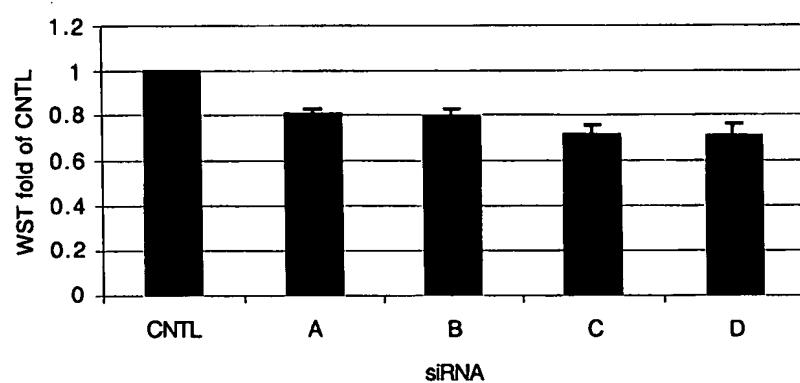


FIGURE 20A

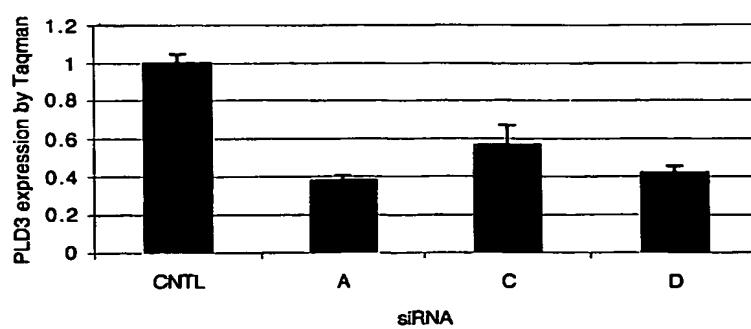


FIGURE 20B

17/24

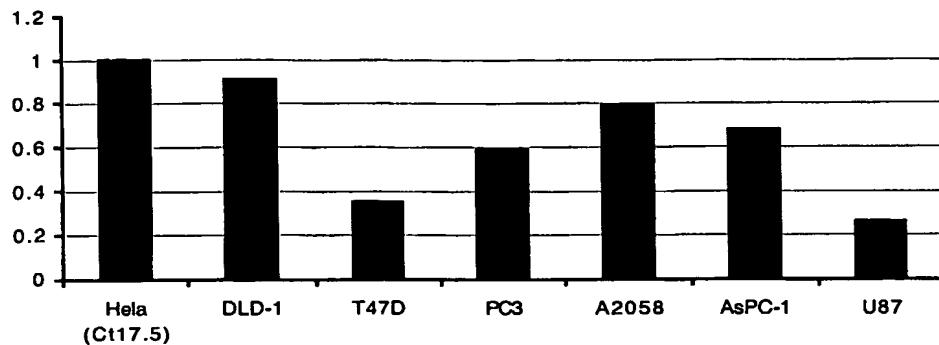


FIGURE 21

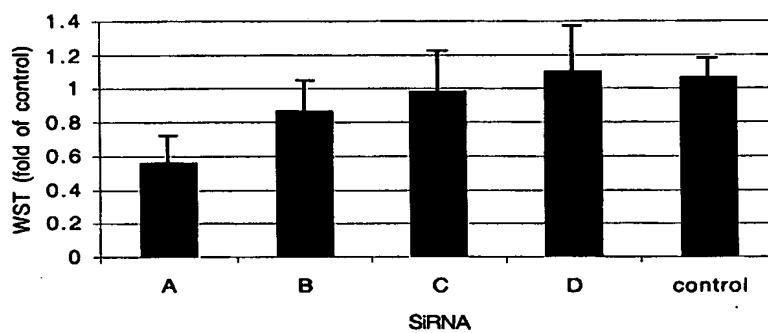


FIGURE 22A

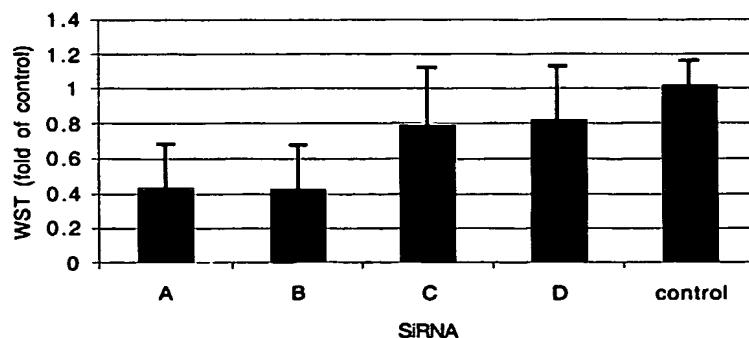


FIGURE 22B

18/24

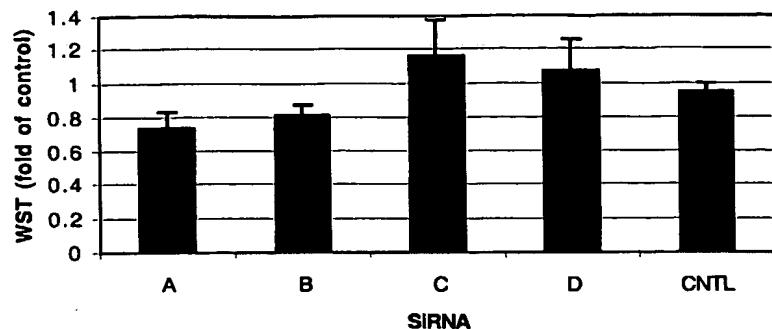


FIGURE 23A

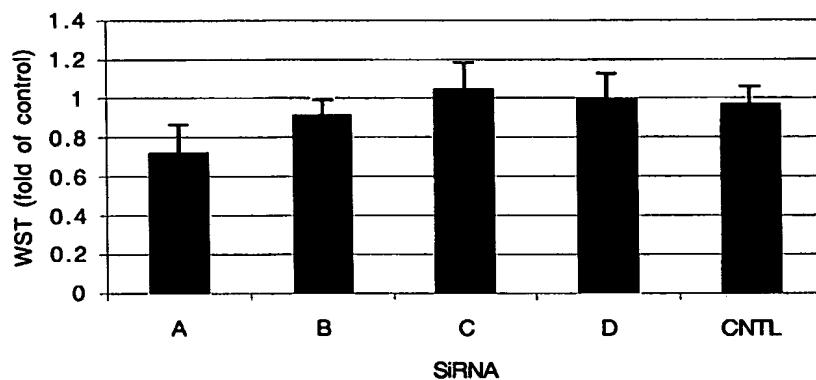


FIGURE 23B

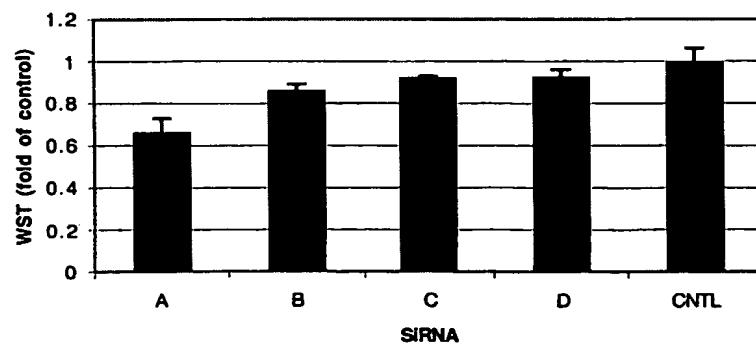


FIGURE 24

19/24

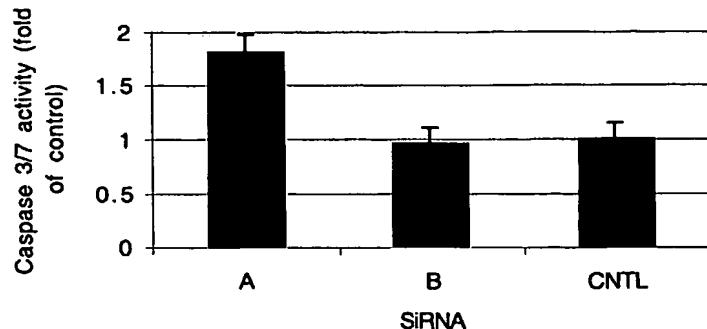


FIGURE 25

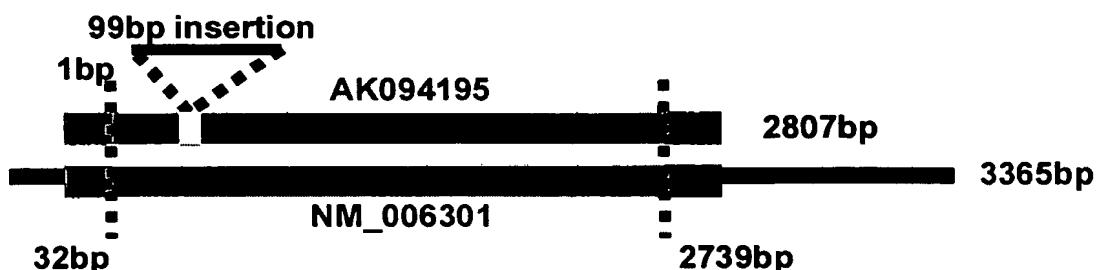


FIGURE 26

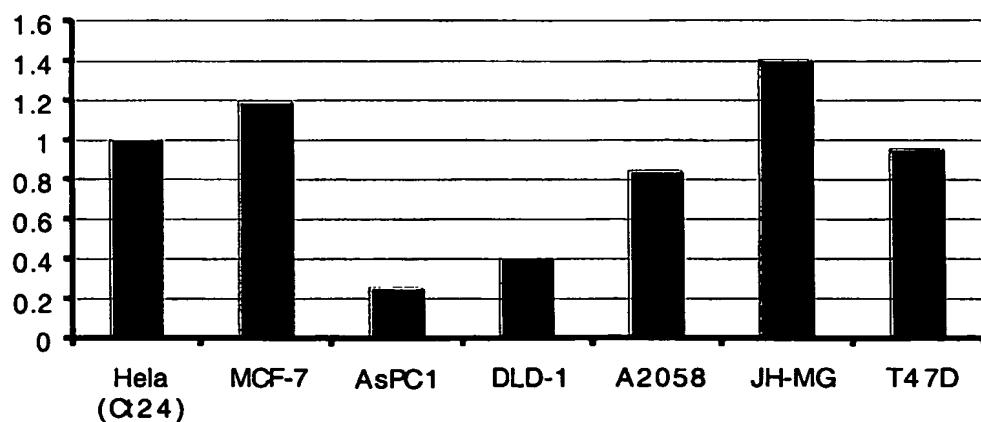


FIGURE 27

20/24



FIGURE 28A

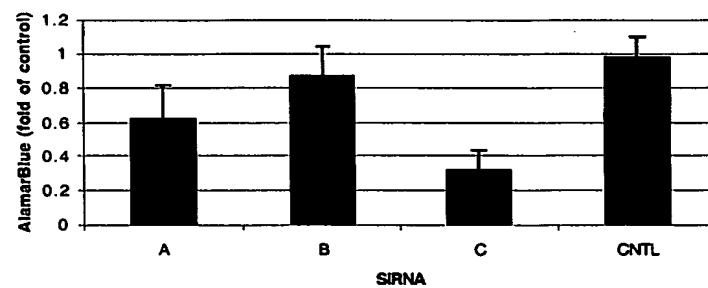


FIGURE 28B

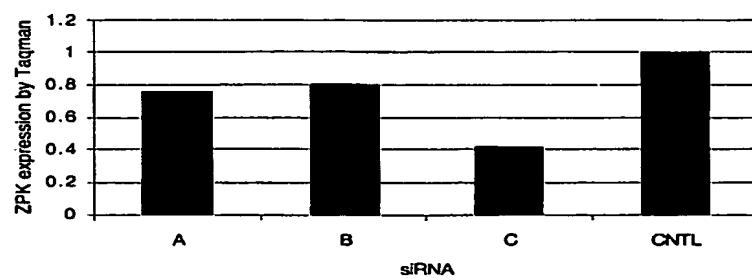


FIGURE 28C

21/24

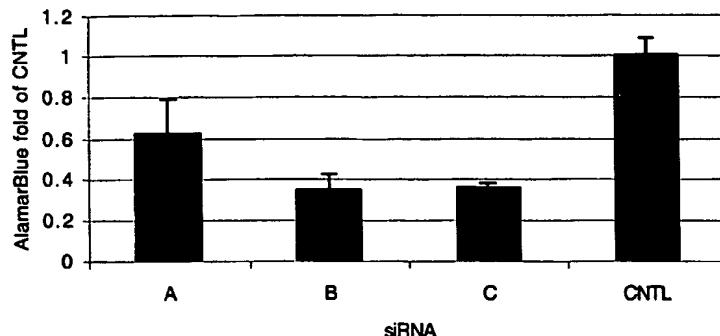


FIGURE 29A

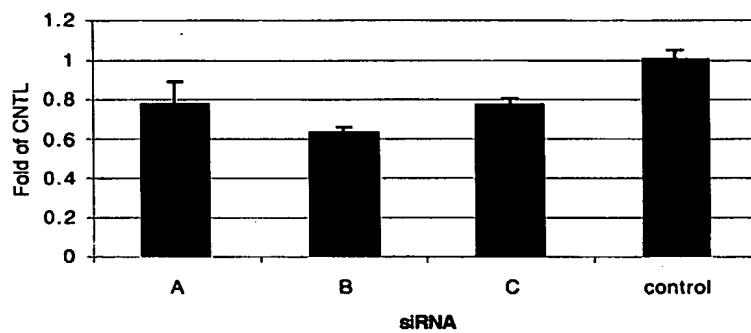


FIGURE 29B

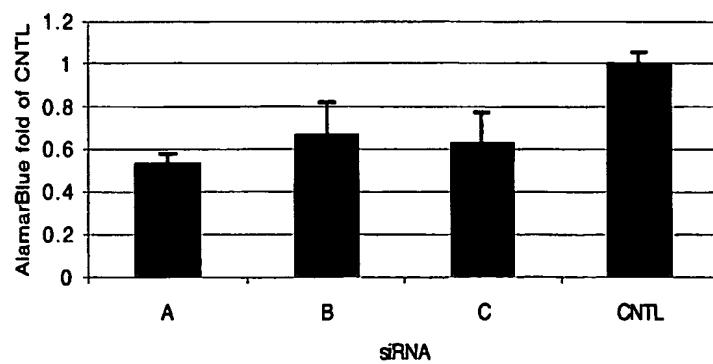


FIGURE 29C

22/24

HCT116

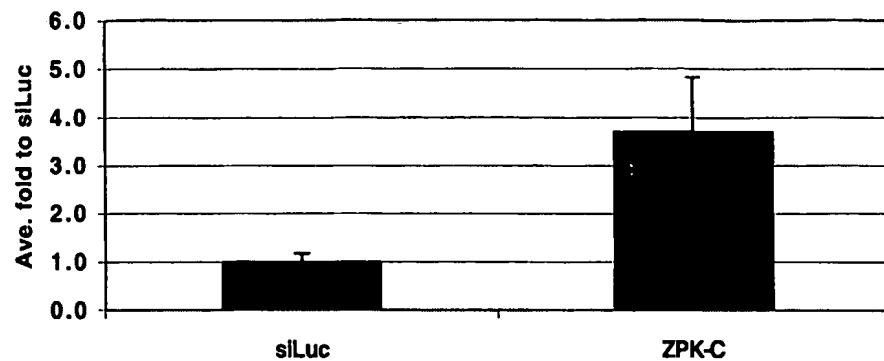


FIGURE 30A

PC3M

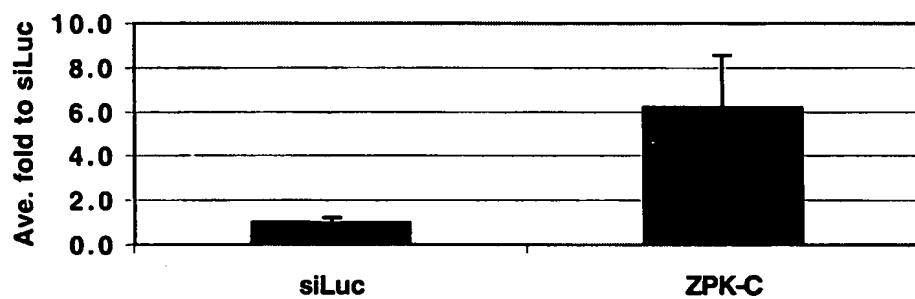


FIGURE 30B

MDAMB231

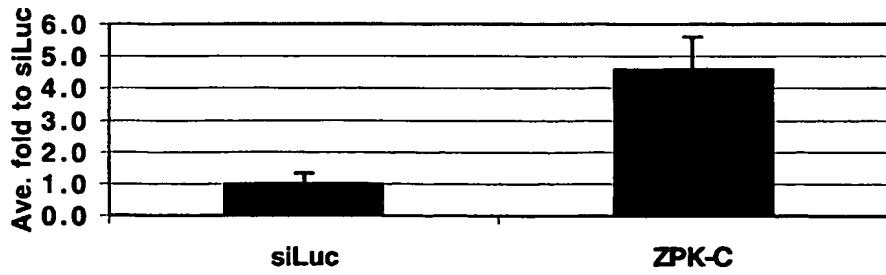


FIGURE 30C

## ZPK Variant 1 DNA Sequence (SEQ ID NO:13)

cttttgtgct	gcggccgcgg	agcccccgag	ggcccaagtgt	tcaccatcat	accaggggcc	60
agaggcgatg	gcttgctcc	atgagacccg	aacaccctct	ccttccttgc	ggggctttgt	120
gtctacccta	agtgaggcat	ccatgcgcaa	gctgacccca	gacacttctg	actgcactcc	180
cgagaaggac	ctgaccccta	cccagtgtgt	acttcgagat	gtggtacccc	tttgtggca	240
gggtggggga	gggcccagcc	cctccccagg	tggagagccg	ccccctgagc	ccttgccaa	300
cagtgtcctg	cagctacatg	agcaggatgc	agggggccca	gggggagcag	ctgggtcacc	360
ttagagtcgg	gcatccagag	ttcgagctga	cgaggtgcga	ctgcagtgcc	agagtggcag	420
tggcttcctt	gaggcctct	ttggctgctc	gcccctgtc	tggaccatga	ttgccaagc	480
ctactccact	gagcacaagc	agcagcagga	agaccttgg	gagggtccct	ttgagggaaat	540
cctggacctg	cagtgggtgg	gctcaggggc	ccaggggtgt	gtcttcctgg	ggcgttcca	600
cggggaggag	gtggctgtga	agaaggtgcg	agacctcaaa	gaaaccgaca	tcaagcactt	660
gcgaaagctg	aagcacccca	acatcatcac	tttcaagggt	gtgtgcaccc	aggctccctg	720
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ctacgacgat	gtggtaaga	tctcagattt	tggcaattcc	aaggagctga	gtgacaagag	960
caccaagatg	tcctttgcag	ggacagtagc	ctggatggcc	cctgaggtga	tcccaatga	1020
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cagtctccat	ctgcccgtgc	cctccagttg	cccagatggt	ttcaagatcc	tgcttcgcca	1200
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cagggagcac	tatgaaagga	agctggagag	agccaaacaac	ctgtatatgg	aacttaatgc	1500
cctcatgttg	cagctggAAC	tcaaggagag	ggagctgtc	aggcgagagc	aagctttaga	1560
gcggaggtgc	ccaggcctgc	tgaagccaca	ccctcccccgg	ggcctctgc	atggaaacac	1620
aatggagaag	cttatacaaga	agaggaatgt	gccacacaa	ctgtcacccc	atagaaaaag	1680
gccagatatc	ctcaagacgg	agtctttgtc	ccctaaacta	gatgcagccc	tgagtgggt	1740
ggggcttcct	gggtgtccta	agggcccccc	ctcaccagga	cgagtcgccc	gtggcaagac	1800
ccgtcaccgc	aaggccagcg	ccaaggggag	ctgtggggac	ctgcctggc	ttctgtacagc	1860
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aatgttctca	tcgtccccag	acctgtgtc	agcagcacta	gggtccccgg	gccggggggc	2040
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tgaacccatc	cccagtggca	cacctgaagt	tggcagcacc	aacactgtat	agcggccaga	2520
tgagcggtct	gatgacatgt	gtctccagg	ctcagaaatc	ccactggacc	caccccttc	2580
agaggtcata	cctggccctg	aaccagctc	cctgcccatt	ccacaccagg	aacttctcag	2640
agagcggggc	cctcccaatt	ctgaggactc	agactgtgac	agcactgaat	tggacaactc	2700
caacagcggt	gatgccttgc	ggccccccagc	ttccctccct	ccatgaaagc	cactcgtatt	2760
ccttgcata	agagaaat	ttatataat	tacatat			2807

FIGURE 31

24/24

**ZPK Variant 1 Amino Acid Sequence (SEQ ID NO:14)**

MACLHETRTP	SPSFGGFVST	LSEASMRKLD	PDTSDCTPEK	DLTPTQCVLR	DVVPLGGQGG	60
GGPSPSPGGE	PPPEPFANSV	LQLHEQDAGG	PGGAAGSPES	RASRVRRADEV	RLQCQSGSGF	120
LEGLFGCLRP	VWTMIGKAYS	TEHKQQQEDL	WEVPFEEILD	LQWVGSGAQQ	AVFLGRFHGE	180
EVAVKKVRDL	KETDIKHLRK	LKHPNIITFK	GVCTQAPCYC	ILMEFCAQQQ	LYEVLRAGRP	240
VTPSLLVDWS	MGIAGGMNYL	HLHKIIHRDL	KSPNMLITYD	DVVKISDFGT	SKELSDKSTK	300
MSFAGTVAWM	APEVIRNEPV	SEKVDIWSFG	VVLWELLTGE	IPYKDVDSSA		350

**FIGURE 32**

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